

```
# https://www.kaggle.com/shayanfazeli/heartbeat # data set

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt # plotting
import os
import sklearn
from keras.models import Sequential
from keras.layers import Flatten, Dense, Dropout, BatchNormalization, AveragePooling2D
from sklearn.model_selection import train_test_split

from sklearn.metrics import accuracy_score, f1_score, precision_score, recall_score
import scipy
from imblearn.over_sampling import SMOTE
from sklearn.model_selection import cross_validate

import tensorflow as tf
from tensorflow import keras
from tensorflow.keras.utils import to_categorical
import numpy as np
import seaborn as sns

# Sample code to verify functionality
print(f"TensorFlow version: {tf.__version__}")
print(f"Keras version: {keras.__version__}")

# Create some sample data
data = np.array([[1, 2, 3], [4, 5, 6]])
labels = np.array([0, 1])

# One-hot encoding of labels
labels_categorical = to_categorical(labels)

print("Labels (categorical):", labels_categorical)

🔗 TensorFlow version: 2.18.0
Keras version: 3.6.0
Labels (categorical): [[1. 0.]
 [0. 1.]]


from google.colab import drive
drive.mount('/content/drive')

import numpy as np
import pandas as pd
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import train_test_split
from sklearn.metrics import accuracy_score, confusion_matrix, classification_report

# Constants
FEATURE_NAMES = ["age", "sex", "SysBP", "DiaBP", "HR", "weightKg", "heightCm", "BMI"]
NUM_FEATURES = len(FEATURE_NAMES)
```


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# Function to read dataset from a CSV file
def read_dataset(file_path):
    # Load the dataset
    data = pd.read_csv(file_path)

    # Select only the relevant feature columns plus the target
    X = data[FEATURE_NAMES].values # Features (specific columns only)
    y = data.iloc[:, -1].values    # Target variable (last column)

    return X, y

# Function to take user input for symptoms
def get_user_input():
    user_input = []
    print("Enter the values for each feature:")
    for feature in FEATURE_NAMES:
        value = float(input(f"{feature.capitalize()}: "))
        user_input.append(value)
    return np.array(user_input).reshape(1, -1)

# Function to display the predicted result for heart disease
def display_prediction_result(prediction):
    result = "has heart disease" if prediction == 1 else "does not have heart disease"
    print(f"The predicted result for heart disease is: {result}")

# Main function
def main():
    file_path = "/content/Health_heart_experimental.csv" # Replace with your file path
    X, y = read_dataset(file_path)

    # Split the data into training and testing sets
    X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2, random_state=42)

    # Initialize the Random Forest Classifier
    rf_model = RandomForestClassifier(n_estimators=100, random_state=42)

    # Train the model with the correct number of features
    rf_model.fit(X_train, y_train)

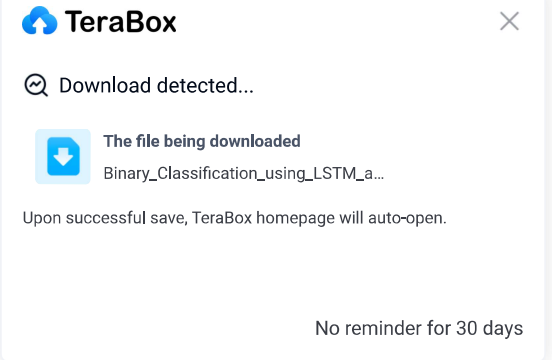
    # Evaluate the model on the test set
    y_pred = rf_model.predict(X_test)
    accuracy = accuracy_score(y_test, y_pred)
    print(f"Model Accuracy: {accuracy:.2f}")
    print("Confusion Matrix:\n", confusion_matrix(y_test, y_pred))
    print("Classification Report:\n", classification_report(y_test, y_pred))

    # Take user input for symptoms
    user_input = get_user_input()

    # Predict the presence of heart disease based on user input
    prediction = rf_model.predict(user_input)[0]

    # Display the predicted result for heart disease
    display_prediction_result(prediction)

```



```
if __name__ == "__main__":
    main()
```

```

Model Accuracy: 0.99
Confusion Matrix:
[[7538  53]
 [ 120 6641]]
Classification Report:

```

	precision	recall	f1-score	support
0	0.98	0.99	0.99	7591
1	0.99	0.98	0.99	6761
accuracy			0.99	14352
macro avg	0.99	0.99	0.99	14352
weighted avg	0.99	0.99	0.99	14352

Enter the values for each feature:

```

import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import random

# Read the training dataset
train = pd.read_csv("/content/drive/MyDrive/archive (2)/mitbih_train.csv", header=None)

# Get the number of samples in the dataset
num_samples = train.shape[0]

# Choose a random sample index
random_index = 90


# Get the signal data for the chosen sample (columns 0 to 186 contain the signal data)
signal_data = train.iloc[random_index, 0:187].values


# Create a time axis (assuming each point is sampled at a fixed interval)
time_axis = np.arange(len(signal_data))


# Get the label of the chosen sample
label = train.iloc[random_index, 187]

# Map the label to its corresponding class
label_map = {0.0: 'N', 1.0: 'S', 2.0: 'V', 3.0: 'F', 4.0: 'Q'}
label_text = label_map.get(label, 'Unknown')

# Plot the signal
plt.figure(figsize=(12, 6))
plt.plot(time_axis, signal_data, 'b-')
plt.title(f'ECG Signal (Class: {label_text})')
plt.xlabel('Sample Index')
plt.ylabel('Amplitude')
plt.grid()
plt.show()
```


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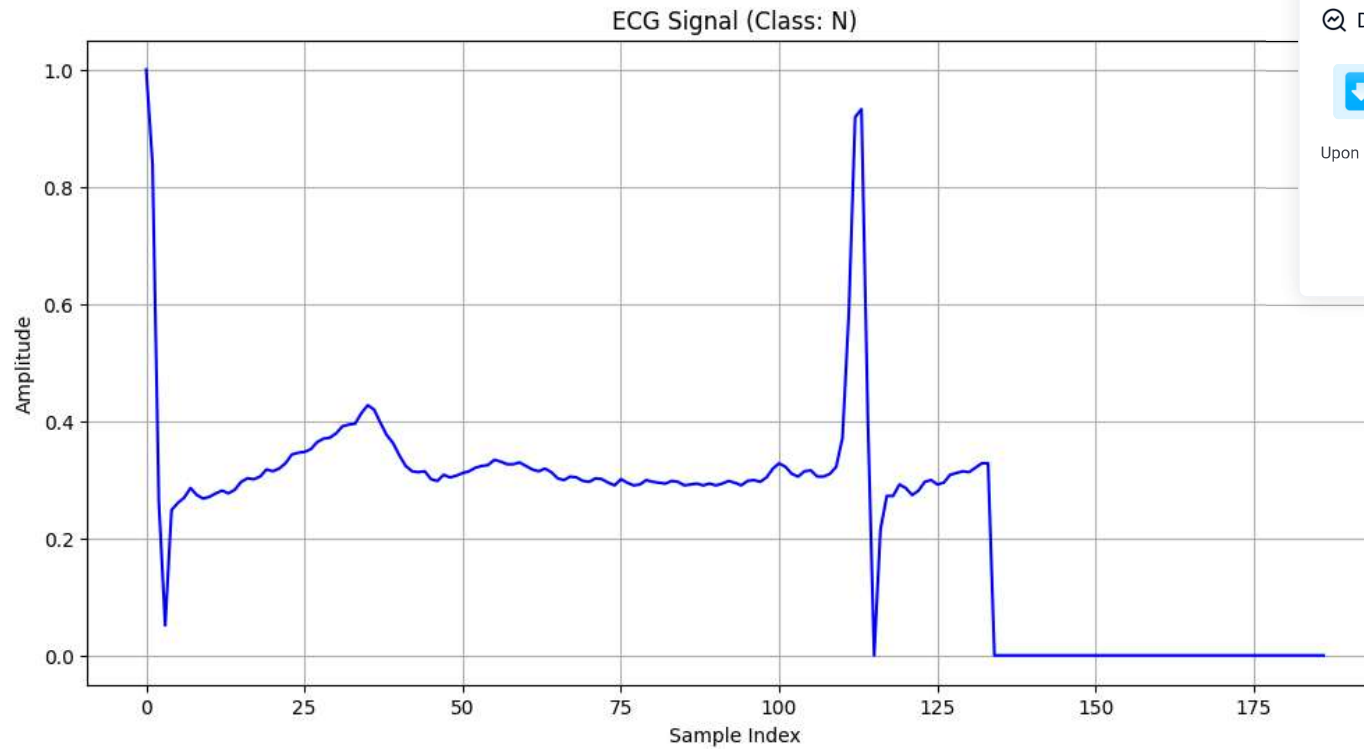


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```
print(f"Displayed signal is from sample index {random_index} with label {label_text}")
```



```
# Assuming num_samples is a list of labels  
num_samples = ["N", "S", "V", "F", "Q", "N", "S", "N", "V"] # Example list
```

```
signal_indices = []  
for i, label in enumerate(num_samples):  
    if label in ["S", "V", "F", "Q"]:  
        signal_indices.append(i)
```

```
print(signal_indices)
```



```
[1, 2, 3, 4, 6, 8]
```

```
# reading csv file
```

```
test=pd.read_csv("/content/drive/MyDrive/archive (2)/mitbih_test.csv",header=None)  
train=pd.read_csv("/content/drive/MyDrive/archive (2)/mitbih_train.csv",header=None)  
#Classes: ['N': 0, 'S': 1, 'V': 2, 'F': 3, 'Q': 4]  
# N- normal, S-supraventricular, V-ventricular, F-fusion, Q- unknown
```

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```
print("Type\tCount")
print((train[187]).value_counts())
```

```
print("train shape : ",train.shape)
print("test shape : ",test.shape)
```

```
↕ Type      Count
187
0.0      72471
4.0       6431
2.0       5788
1.0       2223
3.0        641
Name: count, dtype: int64
train shape : (87554, 188)
test shape  : (21892, 188)
```

```
label_train=[]                                # making multiclass to binary class now labels_train containing either 0 or 1
for i in train.iloc[:,187] :
    if i not in [0.0]:
        label_train.append(1)
        # print("value is",i)
    else:
        label_train.append(0)
        # print("value",i)
```

```
label_test=[]                                # making multiclass to binary class.now labels_test containing either 0 or 1
for i in test.iloc[:,187]:
    if i not in [0.0]:
        label_test.append(1)
        # print("value is",i)
    else:
        label_test.append(0)
        #print("value",i)
```


```
#create new df
labels_train = pd.DataFrame({'col':label_train})
print (labels_train)
```

```
labels_test = pd.DataFrame({'col':label_test})
print (labels_test)
```

```
↕      col
0      0
1      0
2      0
3      0
4      0
...    ...
87549   1
87550   1
```

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```
87551 1
87552 1
87553 1
```

```
[87554 rows x 1 columns]
```

```
col
0 0
1 0
2 0
3 0
4 0
...
21887 1
21888 1
21889 1
21890 1
21891 1
```

```
[21892 rows x 1 columns]
```

```
label_train=np.asarray(label_train)
```

```
label_test=np.asarray(label_test)
```

```
train[187] = labels_train # replace labels column with binary labels
test[187] = labels_test
```

```
print('Count of all classes in training dataset')
print("Type\tCount")
print((train[187]).value_counts())
```

```
print('*****')
```

```
print('Count of all classes in test dataset')
print("Type\tCount")
print((test[187]).value_counts())
```

```
↔ Count of all classes in training dataset
```

```
Type    Count
```

```
187
```

```
0    72471
```

```
1    15083
```

```
Name: count, dtype: int64
```

```
*****
```

```
Count of all classes in test dataset
```

```
Type    Count
```

```
187
```

```
0    18118
```

```
1     3774
```

```
Name: count, dtype: int64
```

```
print("train shape : ",train.shape)
```

```
print("test shape : ",test.shape)
```

```
feature_train=train.iloc[:,0:187]
```

```
#feature_train=train.iloc[:,3]
```

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```
print("feature_train shape : ",feature_train.shape)
```

```
labels_train=train.iloc[:,187]  
print("labels_train shape : ",labels_train.shape)
```

```
feature_test=test.iloc[:,0:187]  
#feature_test=test.iloc[:,3]  
print("feature_test shape : ",feature_test.shape)
```

```
labels_test=test.iloc[:,187]  
print("labels_test shape : ",labels_test.shape)
```

```
↔ train shape : (87554, 188)  
test shape : (21892, 188)  
feature_train shape : (87554, 187)  
labels_train shape : (87554,)  
feature_test shape : (21892, 187)  
labels_test shape : (21892,)
```

```
from sklearn.preprocessing import StandardScaler  
standardized_training_data=StandardScaler().fit_transform(feature_train)  
feature_train=standardized_training_data  
print(feature_train.shape)
```

```
standardized_test_data=StandardScaler().fit_transform(feature_test)  
feature_test=standardized_test_data  
print(feature_test.shape)
```

```
↔ (87554, 187)  
(21892, 187)
```

```
!pip install np_utils  
!pip install keras --upgrade  
!pip install keras
```

```
↔ Requirement already satisfied: np_utils in /usr/local/lib/python3.10/dist-packages (0.6.0)  
Requirement already satisfied: numpy>=1.0 in /usr/local/lib/python3.10/dist-packages (from np_utils) (1.26.4)  
Requirement already satisfied: keras in /usr/local/lib/python3.10/dist-packages (3.6.0)  
Requirement already satisfied: absl-py in /usr/local/lib/python3.10/dist-packages (from keras) (1.4.0)  
Requirement already satisfied: numpy in /usr/local/lib/python3.10/dist-packages (from keras) (1.26.4)  
Requirement already satisfied: rich in /usr/local/lib/python3.10/dist-packages (from keras) (13.9.3)  
Requirement already satisfied: namex in /usr/local/lib/python3.10/dist-packages (from keras) (0.0.8)  
Requirement already satisfied: h5py in /usr/local/lib/python3.10/dist-packages (from keras) (3.11.0)  
Requirement already satisfied: optree in /usr/local/lib/python3.10/dist-packages (from keras) (0.13.0)  
Requirement already satisfied: ml-dtypes in /usr/local/lib/python3.10/dist-packages (from keras) (0.4.1)  
Requirement already satisfied: packaging in /usr/local/lib/python3.10/dist-packages (from keras) (24.1)  
Requirement already satisfied: typing-extensions>=4.5.0 in /usr/local/lib/python3.10/dist-packages (from optree->keras) (4.12.2)  
Requirement already satisfied: markdown-it-py>=2.2.0 in /usr/local/lib/python3.10/dist-packages (from rich->keras) (3.0.0)  
Requirement already satisfied: pygments<3.0.0,>=2.13.0 in /usr/local/lib/python3.10/dist-packages (from rich->keras) (2.18.0)  
Requirement already satisfied: mdurl~=0.1 in /usr/local/lib/python3.10/dist-packages (from markdown-it-py>=2.2.0->rich->keras) (0.1.2)  
Requirement already satisfied: keras in /usr/local/lib/python3.10/dist-packages (3.6.0)  
Requirement already satisfied: absl-py in /usr/local/lib/python3.10/dist-packages (from keras) (1.4.0)  
Requirement already satisfied: numpy in /usr/local/lib/python3.10/dist-packages (from keras) (1.26.4)
```



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```
Requirement already satisfied: rich in /usr/local/lib/python3.10/dist-packages (from keras) (13.9.3)
Requirement already satisfied: namex in /usr/local/lib/python3.10/dist-packages (from keras) (0.0.8)
Requirement already satisfied: h5py in /usr/local/lib/python3.10/dist-packages (from keras) (3.11.0)
Requirement already satisfied: optree in /usr/local/lib/python3.10/dist-packages (from keras) (0.13.0)
Requirement already satisfied: ml-dtypes in /usr/local/lib/python3.10/dist-packages (from keras) (0.4.1)
Requirement already satisfied: packaging in /usr/local/lib/python3.10/dist-packages (from keras) (24.1)
Requirement already satisfied: typing-extensions>=4.5.0 in /usr/local/lib/python3.10/dist-packages (from optree->keras) (4.11.0)
Requirement already satisfied: markdown-it-py>=2.2.0 in /usr/local/lib/python3.10/dist-packages (from rich->keras) (3.0.0)
Requirement already satisfied: pygments<3.0.0,>=2.13.0 in /usr/local/lib/python3.10/dist-packages (from rich->keras) (2.18.0)
Requirement already satisfied: mdurl~=0.1 in /usr/local/lib/python3.10/dist-packages (from markdown-it-py>=2.2.0->rich->keras) (0.1.2)
```

```
x_train1=feature_train
x_test1=feature_test
y_train1=labels_train
y_test1=labels_test
```

```
# Add an Extra Dimension
x_train1=x_train1.reshape(x_train1.shape[0],1,x_train1.shape[1])
x_test1=x_test1.reshape(x_test1.shape[0],1,x_test1.shape[1])
```

```
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras.utils import to_categorical
import numpy as np
import seaborn as sns
```

```
# Sample code to verify functionality
print(f"TensorFlow version: {tf.__version__}")
print(f"Keras version: {keras.__version__}")
```


```
# Create some sample data
data = np.array([[1, 2, 3], [4, 5, 6]])
labels = np.array([0, 1])
```


```
# One-hot encoding of labels
labels_categorical = to_categorical(labels)
```


```
print("Labels (categorical):", labels_categorical)
```

```
TensorFlow version: 2.18.0
Keras version: 3.6.0
Labels (categorical): [[1. 0.]
 [0. 1.]]
```

```
import numpy as np
import tensorflow.keras as keras
from tensorflow.keras.utils import to_categorical
import seaborn as sns
from tensorflow.keras.initializers import RandomNormal
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Dense, LSTM, Embedding
from tensorflow.keras.preprocessing import sequence
```


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```
# fixed random seed for reproducibility
np.random.seed(7)

# here we are having a class number for each sample
print("Class label of first sample:", y_train1[83456])

# lets convert this into a 10 dimensional vector
# ex: consider an image is 5 convert it into 5 => [0, 0, 0, 0, 0, 1, 0, 0, 0, 0]
Y_train = to_categorical(label_train, 2)
Y_test = to_categorical(label_test, 2)
Y_train = to_categorical(y_train1)
print("After converting the output into a vector : ", Y_train[83456])
Y_test = to_categorical(y_test1)
```

```
↗ Class label of first sample: 1
  After converting the output into a vector : [0. 1.]
```

```
print("feature_train",x_train1.shape)
print("Y_train",Y_train.shape)
print("feature_test",x_test1.shape)
print("Y_test",Y_test.shape)
```

```
↗ feature_train (87554, 1, 187)
  Y_train (87554, 2)
  feature_test (21892, 1, 187)
  Y_test (21892, 2)
```

```
output_dim = 2
input_dim = feature_train.shape[1]
```

```
batch_size = 512
nb_epoch = 100
```

```
model_lstm = Sequential()
#model_lstm.add(Embedding)
model_lstm.add(LSTM(100))
model_lstm.add(Dense(output_dim, input_dim=input_dim, activation='softmax'))
```

```
#model_lstm.summary()
```

```
model_lstm.compile(loss='binary_crossentropy', optimizer='adam',metrics=['accuracy'])
```

```
↗ /usr/local/lib/python3.10/dist-packages/keras/src/layers/core/dense.py:87: UserWarning: Do not pass an `input_shape`/`input_dim` argument to a layer. When using Sequential mod
  super().__init__(activity_regularizer=activity_regularizer, **kwargs)
```

```
model_lstm.fit(x_train1, Y_train, epochs=100,batch_size=512)
```

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Epoch 1/100
172/172 6s 12ms/step - accuracy: 0.8576 - loss: 0.3691
Epoch 2/100
172/172 3s 13ms/step - accuracy: 0.9517 - loss: 0.1545
Epoch 3/100
172/172 2s 14ms/step - accuracy: 0.9610 - loss: 0.1280
Epoch 4/100
172/172 3s 14ms/step - accuracy: 0.9651 - loss: 0.1153
Epoch 5/100
172/172 4s 23ms/step - accuracy: 0.9710 - loss: 0.0932
Epoch 6/100
172/172 3s 18ms/step - accuracy: 0.9721 - loss: 0.0882
Epoch 7/100
172/172 4s 13ms/step - accuracy: 0.9746 - loss: 0.0803
Epoch 8/100
172/172 2s 14ms/step - accuracy: 0.9768 - loss: 0.0742
Epoch 9/100
172/172 2s 13ms/step - accuracy: 0.9781 - loss: 0.0688
Epoch 10/100
172/172 4s 22ms/step - accuracy: 0.9804 - loss: 0.0631
Epoch 11/100
172/172 3s 18ms/step - accuracy: 0.9818 - loss: 0.0600
Epoch 12/100
172/172 4s 13ms/step - accuracy: 0.9831 - loss: 0.0541
Epoch 13/100
172/172 3s 14ms/step - accuracy: 0.9838 - loss: 0.0536
Epoch 14/100
172/172 2s 13ms/step - accuracy: 0.9834 - loss: 0.0546
Epoch 15/100
172/172 4s 23ms/step - accuracy: 0.9777 - loss: 0.0670
Epoch 16/100
172/172 4s 14ms/step - accuracy: 0.9856 - loss: 0.0467
Epoch 17/100
172/172 2s 13ms/step - accuracy: 0.9855 - loss: 0.0460
Epoch 18/100
172/172 2s 12ms/step - accuracy: 0.9867 - loss: 0.0433
Epoch 19/100
172/172 2s 14ms/step - accuracy: 0.9872 - loss: 0.0415
Epoch 20/100
172/172 4s 22ms/step - accuracy: 0.9872 - loss: 0.0397
Epoch 21/100
172/172 3s 20ms/step - accuracy: 0.9882 - loss: 0.0373
Epoch 22/100
172/172 2s 14ms/step - accuracy: 0.9885 - loss: 0.0371
Epoch 23/100
172/172 2s 13ms/step - accuracy: 0.9884 - loss: 0.0369
Epoch 24/100
172/172 3s 18ms/step - accuracy: 0.9899 - loss: 0.0333
Epoch 25/100
172/172 4s 25ms/step - accuracy: 0.9904 - loss: 0.0324
Epoch 26/100
172/172 4s 25ms/step - accuracy: 0.9907 - loss: 0.0308
Epoch 27/100
172/172 3s 15ms/step - accuracy: 0.9909 - loss: 0.0298
Epoch 28/100
172/172 2s 13ms/step - accuracy: 0.9893 - loss: 0.0338
Epoch 29/100
172/172 2s 13ms/step - accuracy: 0.9913 - loss: 0.0285

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```
predictions=model_lstm.predict(x_test1)
```

```
↔ 685/685 ————— 2s 3ms/step
```

```
print(classification_report(Y_test.argmax(axis=1), predictions.argmax(axis=1)))
```

```
↔
```

	precision	recall	f1-score	support
0	0.99	0.99	0.99	18118
1	0.96	0.94	0.95	3774
accuracy			0.98	21892
macro avg	0.97	0.97	0.97	21892
weighted avg	0.98	0.98	0.98	21892

```
print(predictions)
```

```
a=predictions.argmax(axis=1) # 0 is row and 1 is column
```

```
#print(a)
```

```
#create new df
```

```
a = pd.DataFrame({'col':a})
```

```
print("Type\tCount")
```

```
print((a['col']).value_counts())
```

```
↔ [[9.9999994e-01 3.3045269e-15]
 [9.9999994e-01 3.0559510e-10]
 [9.9999994e-01 1.4586851e-10]
 ...
 [6.3516182e-14 1.0000000e+00]
 [8.8528944e-22 1.0000000e+00]
 [4.4833044e-19 1.0000000e+00]]
Type    Count
col
0      18199
1       3693
Name: count, dtype: int64
```

```
#
```

	precision	recall	f1-score
0	0.99	0.99	0.99
1	0.96	0.94	0.95

```
# accuracy : 0.98
```

```
# Evaluate the model
```

```
predictions = model_lstm.predict(x_test1)
```

```
print(classification_report(Y_test.argmax(axis=1), predictions.argmax(axis=1)))
```


```
# Save the model for future use
```

```
model_lstm.save("/content/drive/MyDrive/mit-bh/heartbeat_classifier.h5")
```

```
"Function to preprocess a single EEG signal"
```

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```

# Function to preprocess a single ECG signal
def preprocess_signal(signal):
    # Ensure the signal has the correct length (187 in this case)
    if len(signal) != 187:
        raise ValueError("Signal must have 187 data points")

    # Reshape for sklearn's StandardScaler
    signal = np.array(signal).reshape(1, -1)

    # Use the same scaler that was used for training data
    standardized_signal = StandardScaler().fit_transform(signal)

    # Reshape for LSTM input (samples, timesteps, features)
    return standardized_signal.reshape(1, 1, 187)

# Function to predict heartbeat type
def predict_heartbeat(signal):
    try:
        processed_signal = preprocess_signal(signal)
        prediction = model_lstm.predict(processed_signal)
        class_idx = prediction.argmax(axis=1)[0]

        if class_idx == 0:
            return "Normal Heartbeat"
        else:
            return "Abnormal Heartbeat"
    except Exception as e:
        return f"Error: {str(e)}"

# Example: Let's take a sample from the test set
sample_idx = 9 # You can change this index to try different samples
sample_signal = feature_test[sample_idx]
true_label = "Normal" if label_test[sample_idx] == 0 else "Abnormal"


print(f"Analyzing sample signal {sample_idx}...")
print("True Label:", true_label)
print("Prediction:", predict_heartbeat(sample_signal))


# Let's also visualize this ECG signal
plt.figure(figsize=(12, 4))
plt.plot(sample_signal)
plt.title(f'ECG Signal {sample_idx} (True: {true_label})')
plt.xlabel('Time')
plt.ylabel('Amplitude')
plt.grid()
plt.show()


# Now let's create a custom ECG signal for testing
# This is just a dummy example; in practice, you'd use real ECG data
sampling_rate = 100
data_length = 187
ecg_data = np.random.normal(0, 1, data_length)

# Add some noise to the data
noise_amplitude = 0.5
ecg_data += np.random.normal(0, noise_amplitude, data_length)

```


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```
# Scale the data to be between -1 and 1
ecg_data /= np.max(np.abs(ecg_data))

# sampling_rate = 360
# data_length = 187

# # Generate random data with a normal distribution
# ecg_data = np.random.normal(0, 1, data_length)


# # Add noise to the data
# noise_amplitude = 0.5
# noise = np.random.normal(0, noise_amplitude, data_length)
# noisy_ecg_data = ecg_data + noise


# # Scale the data to be between -1 and 1
# noisy_ecg_data /= np.max(np.abs(noisy_ecg_data))


print("\nAnalyzing custom signal...")
print("Prediction:", predict_heartbeat(ecg_data))


# Visualize the custom signal
plt.figure(figsize=(12, 4))
plt.plot(ecg_data)
plt.title('Custom ECG Signal')
plt.xlabel('Time')
plt.ylabel('Amplitude')
plt.grid()
plt.show()

# You can also test the model with signals from scientific papers or other sources
# Just make sure they have 187 data points or interpolate/truncate them to fit
```

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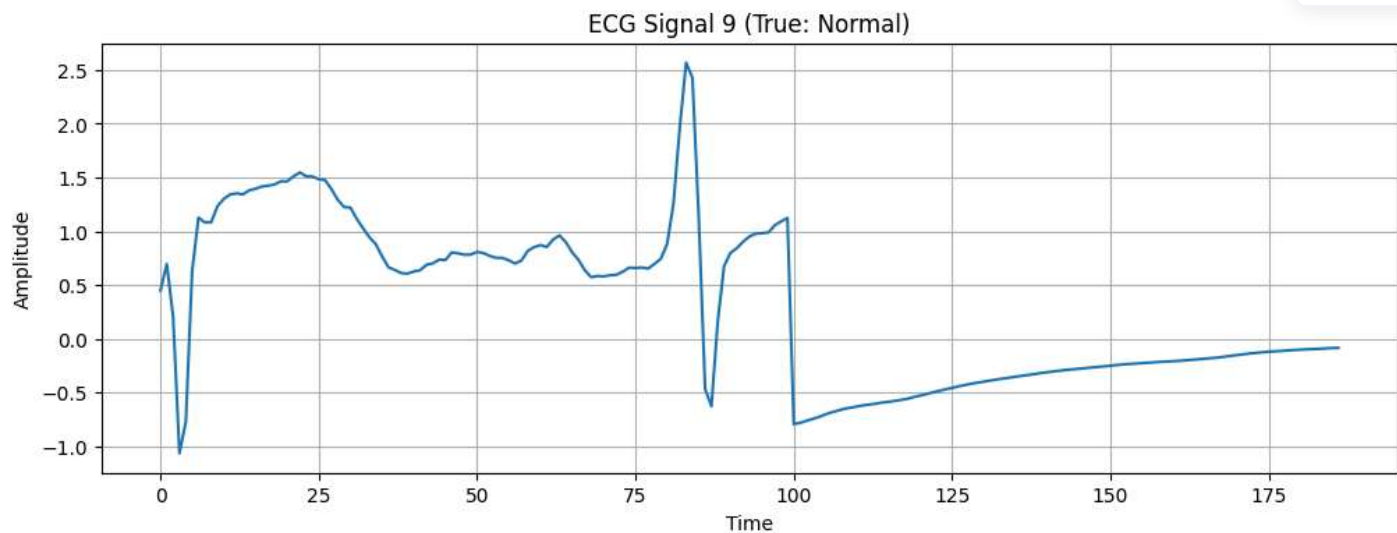


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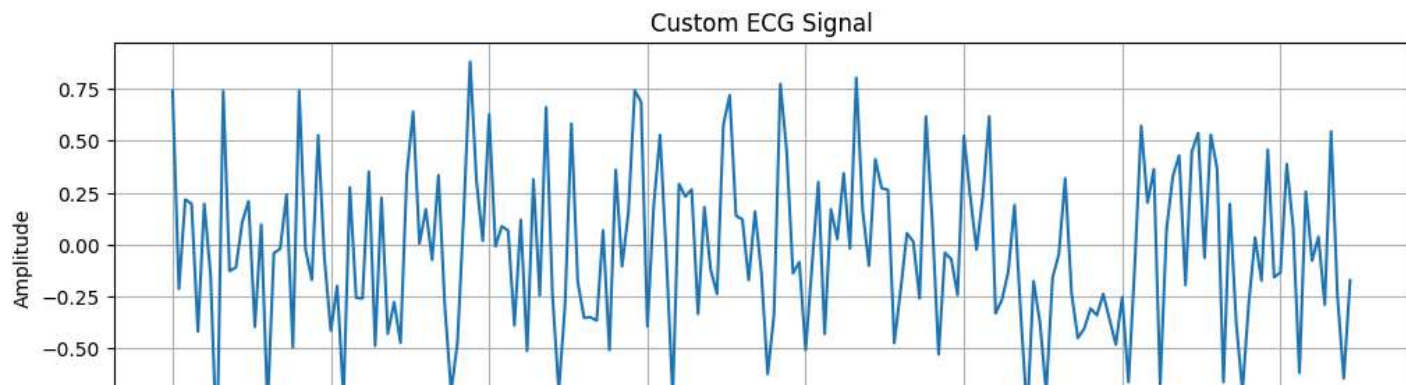
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685/685 2s 2ms/step
WARNING:absl:You are saving your model as an HDF5 file via `model.save()` or `keras.saving.save_model(model)`. This file
precision recall f1-score support
0 0.99 0.99 0.99 18118
1 0.96 0.94 0.95 3774
accuracy 0.98 21892
macro avg 0.97 0.97 0.97 21892
weighted avg 0.98 0.98 0.98 21892
Analyzing sample signal 9...
True Label: Normal
1/1 0s 49ms/step
Prediction: Normal Heartbeat



Analyzing custom signal...
1/1 0s 47ms/step
Prediction: Normal Heartbeat



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